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Amendments t Claims

This listing of claims will replace all prior versions and listings of claims in the application:

Claim 1. (currently amended) An isolated nucleic acid sequence encoding a polypeptide with isoflavone synthase activity having the amino acid sequence set forth in SEQ ID NO:66, wherein the nucleic acid fragment does not have the nucleotide sequence set forth in SEQ ID NO:9, and wherein

Xaa₁₀ is Phe or Leu

Xaa₁₆ is Ser or Leu

Xaa₂₃ is Ser or Thr

Xaa₂₅ is lle or Lys

Xaa₃₉ is Lys or Arg

Xaa₄₈ is Pro or Leu

Xaa₆₀ is Pro or Leu

Xaa₇₃ is Leu or His

Xaa₇₄ is Ser or Tyr

Xaa₉₅ is Ala or Thr

Xaa₉₆ is Asn or His

Xaa₁₀₂ is Asn or Ser

Xaa₁₁₀ is Ile, Val, or Thr

Xaa₁₁₂ is Arg or His

Xaa₁₁₇ is Asn or Ser

Xaa₁₁₈ is Ser or Leu

Xaa₁₂₁ is Met or Arg

Xaa₁₂₂ is Ala or Val

Xaa₁₂₄ is Phe or Ile

Xaa₁₂₉ is Lys or Arg

Xaa₁₄₇ is Lys or Glu

Xaa₁₅₉ is Leu or Phe

Xaa₁₆₂ is Ala or Val

Xaa₁₆₆ is Ser or Gly

Xaa₁₇₀ is Gln or Arg

Xaa₁₇₅ is Val or Leu

Xaa₁₈₃ is Ala or Thr

Xaa₁₈₇ is Thr or lle

Xaa₁₉₁ is Met or Val

Xaa₂₀₉ is Phe or Tyr

Xaa₂₁₉ is Arg or Trp

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Xaa₂₂₃ is Tyr or His

Xaa₂₅₃ is Gly or Glu

Xaa₂₅₉ is Lys or Glu

Xaa₂₆₃ is Val or Asp

Xaa₂₆₄ is Val, Asp, or lle

Xaa₂₆₈ is Ala or Val

Xaa₂₇₂ is Phe or Leu

Xaa₂₈₅ is Thr or Met

Xaa₂₉₃₂₉₂ is Glu or Asp

Xaa₂₉₃ is GIn or His

Xaa294 is Thr, or Ile

Xaa₃₀₁ is Phe or Leu

Xaa₃₀₆ is Thr or Ile

Xaa₃₁₁ is Val or Glu

Xaa₃₁₂ is Val or Ala

Xaa₃₂₅ is Arg or Lys

Xaa₃₂₈ is Gln or Glu

Xaa₃₃₄ is Val or Ala

Xaa₃₄₂ is Arg or Ile

Xaa₃₇₇ is Thr or lle

Xaa381 is Glu or Gly

Xaa₃₈₅ is Tyr, His, or Cys

Xaa₃₈₇ is Ile or Thr

Xaa₃₉₃ is Val or lle

Xaa₃₉₄ is Leu or Pro

Xaa₄₀₂ is Arg or Lys

Xaa₄₀₄ is Ser or Pro

Xaa₄₁₃ is Ser or Phe

Xaa₄₂₂ is Glu or Gly

Xaa₄₂₈ is Gly or Arg

Xaa₄₂₉ is Pro or Leu

Xaa₄₃₅ is Gln or Arg

Xaa₄₄₇ is Arg or Gly

Xaa₄₅₃ is Asn, Ser, or lle

Xaa₄₅₉ is Met or Thr, and

Xaa₄₈₅ is Asp or Gly.

Claim 2. (canceled) An isolated polypeptide sequence of SEQ ID NO: 66 wherein Xaa₁₀ is Phe or Leu

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Xaa₁₆ is Ser or Leu

Xaa₂₃ is Ser or Thr

Xaa₂₅ is lle or Lys

Xaa₃₉ is Lys or Arg

Xaa₄₈ is Pro or Leu

Xaa₆₀ is Pro or Leu

Xaa₇₃ is Leu or His

Xaa₇₄ is Ser or Tyr

Xaa₉₅ is Ala or Thr

Xaa₉₆ is Asn or His

Xaa₁₀₂ is Asn or Ser

Xaa₁₁₀ is Ile, Val, or Thr

Xaa₁₁₂ is Arg or His

Xaa₁₁₇ is Asn or Ser

Xaa₁₁₈ is Ser or leu

Xaa₁₂₁ is Met or Arg

Xaa₁₂₂ is Ala or Val

Xaa₁₂₄ is Phe or Ile

Xaa₁₂₉ is Lys or Arg

Xaa₁₄₇ is Lys or Glu

Xaa₁₅₉ is Leu or Phe

Xaa₁₆₂ is Ala or Val

Xaa₁₆₆ is Ser or Gly

Xaa₁₇₀ is Gln or Arg

Xaa₁₇₅ is Val or Leu

Xaa₁₈₃ is Ala or Thr

Xaa₁₈₇ is Thr or lle

Xaa₁₉₁ is Met or Val

Xaa₂₀₉ is Phe or Tyr

Xaa₂₁₉ is Arg or Trp

Xaa₂₂₃ is Tyr or His

Xaa₂₅₃ is Gly or Glu

Xaa₂₅₉ is Lys or Glu

Xaa₂₆₃ is Val or Asp

Xaa₂₆₄ is Val, Asp, or lle

Xaa₂₆₈ is Ala or Val

Xaa₂₇₂ is Phe or Leu

Xaa₂₈₅ is Thr or Met

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Xaa₂₉₃ is Glu or Asp

Xaa₂₉₄ is Thr, or lle

Xaa₃₀₁ is Phe or Leu

Xaa₃₀₆ is Thr or Ile

Xaa₃₁₁ is Val or Glu

Xaa₃₁₂ is Val or Ala

Xaa₃₂₅ is Arg or Lys

Xaa₃₂₈ is Gln or Glu

Xaa₃₃₄ is Val or Ala

Xaa₃₄₂ is Arg or Ile

Xaa₃₇₇ is Thr or Ile

Xaa₃₈₁ is Glu or Gly

Xaa₃₈₅ is Tyr, His, or Cys

Xaa₃₈₇ is lle or Thr

Xaa393 is Val or lle

Xaa₃₉₄ is Leu or Pro

Xaa₄₀₂ is Arg or Lys

Xaa₄₀₄ is Ser or Pro

Xaa₄₁₃ is Ser or Phe

Xaa₄₂₂ is Glu or Gly

Xaa₄₂₈ is Gly or Arg

Xaa₄₂₉ is Pro or Leu

Xaa₄₃₅ is Gln or Arg

Xaa₄₄₇ is Arg or Gly

Xaa₄₅₃ is Asn, Ser, or lle

Xaa₄₅₉ is Met or Thr, and

Xaa₄₈₅ is Asp or Gly.

Claim 3. (currently amended) An isolated nucleic acid sequence encoding a polypeptide with isoflavone synthase activity capable of converting 2S-flavanone into an isoflavonoid.

Claim 4. (currently amended) An isolated nucleic acid sequence encoding a polypeptide with isoflavone synthase activity capable of converting 2S-flavanone into an isoflavonoid wherein the nucleic acid sequence is not the nucleic acid sequence set forth in SEQ ID NO:9.

Claims 5-10 (canceled).

Claim 11. (currently amended) A chimeric <u>polynucleotide</u> sequence comprising the nucleic acid sequence of Claim 1 operably linked to <u>at least one</u> suitable regulatory sequences sequence.

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Claim 12. (currently amended) A transformed host cell comprising the chimeric polynucleotide sequence of Claim 11.

Claim 13. (currently amended) The transformed host cell of Claim 12 further comprising a second chimeric polynucleotide sequence comprising a nucleic acid sequence encoding a polypeptide that regulates expression of at least one enzyme of the phenylpropanoid pathway.

Claim 14. (currently amended) The transformed host cell of Claim 13 wherein the second chimeric polynucleotide sequence comprises a chimera containing a polynucleotide encoding the maize R region, wherein the R region is between polynucleotide the region encoding the maize C1 DNA binding domain and the maize C1 activation domain.

Claim 15. (original) The transformed host cell of Claim 12 wherein the host cell is a eukaryotic cell.

Claim 16. (original) The eukaryotic cell of Claim 13 wherein the cell is a yeast cell.

Claim 17. (original) The eukaryotic cell of Claim 15 wherein the cell is a plant cell.

Claim 18. (original) The plant cell of Claim 17 wherein the cell is a soybean cell.

Claim 19. (original) The plant cell of Claim 17 wherein the cell is a corn cell.

Claims 20-25. (canceled).

Claim 26. (currently amended) A method of altering the level of expression of isoflavone synthase in a host cell comprising:

- (a) transforming a host cell with the chimeric polynucleotide sequence of Claim 11:
- (b) optionally or transforming the host cell with the chimeric polynucleotide of Claim 11 and with a second chimeric polynucleotide sequence comprising a nucleic acid sequence encoding a polypeptide that regulates expression of at least one enzyme of the phenylpropanoid pathway; and
- (be) growing the transformed host cell produced in step (a) or step (b) under conditions that are suitable for expression of the chimeric polynucleotide sequence

wherein expression of the chimeric sequences result polynucleotide results in production of altered levels of isoflavone synthase in the transformed host cell. Claims 27 and 28 (canceled).

Claim 29. (currently amended) The method of Claim 26 or Claim 27 wherein the host cell is a eukaryotic cell.

Claim 30. (currently amended) The method of Claim 26 or Claim 27 wherein the eukaryotic cell is a yeast cell.

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Claim 31. (currently amended) The method of Claim 26 or Claim 27 wherein the eukaryotic cell is a plant cell.

Claim 32. (original) The method of Claim 31 wherein the plant cell is a soybean cell.

Claim 33. (original) The method of Claim 31 wherein the plant cell is a corn cell.

Claims 34-50 (canceled).

Claim 51. (new) The isolated nucleic acid fragment of Claim 1 where

Xaa₁₀ is Phe

Xaa₁₆ is Leu

Xaa₂₃ is Ser

Xaa₂₅ is Lys

Xaa₃₉ is Lys

Xaa₄₈ is Leu

Xaa₆₀ is Leu

Xaa₇₃ is Leu

Xaa₇₄ is Ser

Xaa₉₅ is Thr

Xaa₉₆ is His

Xaa₁₀₂ is Asn

Xaa₁₁₀ is Ile

Xaa₁₁₂ is Arg

Xaa₁₁₇ is Asn

Xaa₁₁₈ is Ser

Xaa₁₂₁ is Met

Xaa₁₂₂ is Val

Xaa₁₂₄ is Phe

Xaa₁₂₉ is Lys

Xaa₁₄₇ is Lys

Xaa₁₅₉ is Phe

Xaa₁₆₂ is Val

Xaa₁₆₆ is Ser

Xaa₁₇₀ is Gln

Xaa₁₇₅ is Val

Xaa₁₈₃ is Thr

Xaa₁₈₇ is Ile

Xaa₁₉₁ is Met

Xaa₂₀₉ is Phe

Xaa₂₁₉ is Trp

Xaa₂₂₃ is Tyr

Xaa₂₅₃ is Glu

Xaa₂₅₉ is Lys

Xaa₂₆₃ is Val

Xaa₂₆₄ is Val

Xaa₂₆₈ is Ala

Xaa₂₇₂ is Phe

Xaa₂₈₅ is Met

Xaa₂₉₂ is Glu

Xaa₂₉₃ is Gln

Xaa₂₉₄ is Ile

Xaa₃₀₁ is Phe

Xaa₃₀₆ is Thr

Xaa₃₁₁ is Val

Xaa₃₁₂ is Ala

Xaa₃₂₅ is Arg

Xaa₃₂₈ is Gln

Xaa₃₃₄ is Val

Xaa₃₄₂ is Arg

Xaa₃₇₇ is Thr

Xaa₃₈₁ is Glu

Xaa₃₈₅ is Tyr

Xaa₃₈₇ is Ile

Xaa₃₉₃ is Val

Xaa394 is Leu

Xaa₄₀₂ is Arg

Xaa₄₀₄ is Pro

Xaa₄₁₃ is Phe

Xaa₄₂₂ is Gly

Xaa₄₂₈ is Gly

Xaa₄₂₉ is Pro

Xaa₄₃₅ is GIn

Xaa₄₄₇ is Arg

Xaa₄₅₃ is Asn

Xaa₄₅₉ is Met, and

Xaa₄₈₅ is Asp.

Claim 52. (new) The isolated nucleic acid fragment of Claim 1 where

Xaa₁₀ is Phe or Leu

Xaa₁₆ is Leu

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Xaa₂₃ is Ser

Xaa₂₅ is Lys

Xaa₃₉ is Lys

Xaa₄₈ is Leu

Xaa₆₀ is Leu

Xaa₇₃ is Leu

Xaa₇₄ is Ser

Xaa₉₅ is Thr

Xaa₉₆ is His

Xaa₁₀₂ is Asn

Xaa₁₁₀ is Thr

Xaa₁₁₂ is Arg

Xaa₁₁₇ is Asn

Xaa₁₁₈ is Ser

Xaa₁₂₁ is Met

Xaa₁₂₂ is Val

Xaa₁₂₄ is Phe

Xaa₁₂₉ is Arg

Xaa₁₄₇ is Lys

Xaa₁₅₉ is Phe

Xaa₁₆₂ is Val

Xaa₁₆₆ is Ser

Xaa₁₇₀ is Gln

Xaa₁₇₅ is Val

Xaa₁₈₃ is Thr

Xaa₁₈₇ is Ile

Xaa₁₉₁ is Met

Xaa₂₀₉ is Phe

Xaa₂₁₉ is Trp

Xaa₂₂₃ is Tyr

Xaa₂₅₃ is Gly

Xaa₂₅₉ is Glu

Xaa₂₆₃ is Val

Xaa₂₆₄ is Val

Xaa₂₆₈ is Ala

Xaa₂₇₂ is Phe

Xaa₂₈₅ is Met

Xaa₂₉₂ is Glu

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Xaa₂₉₃ is Gln

Xaa₂₉₄ is Ile

Xaa₃₀₁ is Leu

Xaa₃₀₆ is Thr

Xaa₃₁₁ is Val

Xaa₃₁₂ is Ala

Xaa₃₂₅ is Arg

Xaa₃₂₈ is Gln

Xaa₃₃₄ is Val

Xaa₃₄₂ is Arg

Xaa₃₇₇ is Thr

Xaa₃₈₁ is Glu

Xaa₃₈₅ is Tyr

303

Xaa₃₈₇ is lle

Xaa₃₉₃ is Val

Xaa₃₉₄ is Leu

Xaa₄₀₂ is Arg

Xaa₄₀₄ is Pro

Xaa₄₁₃ is Phe

Xaa₄₂₂ is Gly

Xaa₄₂₈ is Gly

Xaa₄₂₉ is Pro

Xaa₄₃₅ is Gln

Xaa₄₄₇ is Arg

Xaa₄₅₃ is Asn

Xaa₄₅₉ is Met, and

Xaa₄₈₅ is Asp.

Claim 53. (new) The isolated nucleic acid fragment of Claim 1 where

Xaa₁₀ is Phe or Leu

Xaa₁₆ is Leu

Xaa₂₃ is Thr

Xaa₂₅ is Lys

Xaa₃₉ is Lys

Xaa₄₈ is Leu

Xaa₆₀ is Leu

Xaa₇₃ is Leu

Xaa₇₄ is Tyr

Xaa₉₅ is Thr

Xaa₉₆ is His

Xaa₁₀₂ is Asn

Xaa₁₁₀ is Ile

Xaa₁₁₂ is Arg

Xaa₁₁₇ is Ser

Xaa₁₁₈ is Leu

Xaa₁₂₁ is Met

Xaa₁₂₂ is Val

Xaa₁₂₄ is Phe

Xaa₁₂₉ is Lys

Xaa₁₄₇ is Lys

Xaa₁₅₉ is Phe

Xaa₁₆₂ is Val

Xaa₁₆₆ is Gly

Xaa₁₇₀ is Gln

Xaa₁₇₅ is Leu

Xaa₁₈₃ is Thr

Xaa₁₈₇ is Ile

Xaa₁₉₁ is Met

Xaa₂₀₉ is Tyr

Xaa₂₁₉ is Trp

Xaa₂₂₃ is His

Xaa₂₅₃ is Glu

Xaa₂₅₉ is Lys

Xaa₂₆₃ is Val

Xaa₂₆₄ is Val

Xaa₂₆₈ is Val

Xaa₂₇₂ is Phe

Xaa₂₈₅ is Thr

Xaa₂₉₂ is Asp

Xaa₂₉₃ is His

Xaa₂₉₄ is Ile

Xaa₃₀₁ is Phe

Xaa₃₀₆ is Ile

Xaa₃₁₁ is Val

Xaa₃₁₂ is Ala Xaa₃₂₅ is Lys

Xaa₃₂₈ is Glu

Xaa₃₃₄ is Val

Xaa₃₄₂ is Arg

Xaa₃₇₇ is Thr

Xaa₃₈₁ is Glu

Xaa₃₈₅ is Tyr

Xaa₃₈₇ is Ile

Xaa₃₉₃ is Ile

Xaa₃₉₄ is Leu

Xaa₄₀₂ is Arg

Xaa₄₀₄ is Pro

Xaa₄₁₃ is Phe

Xaa₄₂₂ is Gly

Xaa₄₂₈ is Arg

Xaa₄₂₉ is Pro

Xaa₄₃₅ is Gln

Xaa₄₄₇ is Gly

Xaa₄₅₃ is Asn

Xaa₄₅₉ is Met, and

Xaa₄₈₅ is Gly.

Claim 54. (new) The isolated nucleic acid fragment of Claim 1 where

Xaa₁₀ is Phe or Leu

Xaa₁₆ is Leu

Xaa₂₃ is Thr

Xaa₂₅ is Lys

Xaa₃₉ is Lys

Xaa₄₈ is Pro

Xaa₆₀ is Leu

Xaa₇₃ is Leu

Xaa₇₄ is Tyr

Xaa₉₅ is Thr

Xaa₉₆ is His

Xaa₁₀₂ is Asn

Xaa₁₁₀ is lle

Xaa₁₁₂ is Arg

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Xaa₁₁₇ is Ser

Xaa₁₁₈ is Ser

Xaa₁₂₁ is Met

Xaa₁₂₂ is Val

Xaa₁₂₄ is Phe

Xaa₁₂₉ is Lys

Xaa₁₄₇ is Lys

Xaa₁₅₉ is Phe

Xaa₁₆₂ is Val

Xaa₁₆₆ is Ser

Xaa₁₇₀ is Gln

Xaa₁₇₅ is Val

Xaa₁₈₃ is Thr

Xaa₁₈₇ is lle

Xaa₁₉₁ is Met

Xaa₂₀₉ is Phe

Xaa₂₁₉ is Trp

Xaa₂₂₃ is Tyr

Xaa₂₅₃ is Glu

Xaa₂₅₉ is Lys

Xaa₂₆₃ is Val

Xaa₂₆₄ is Val

Xaa₂₆₈ is Ala

Xaa₂₇₂ is Phe

Xaa₂₈₅ is Met

Xaa₂₉₂ is Glu

Xaa₂₉₃ is Gln

Xaa₂₉₄ is lle

Xaa₃₀₁ is Phe

Xaa₃₀₆ is Thr

Xaa₃₁₁ is Val

Xaa₃₁₂ is Ala

Xaa₃₂₅ is Arg

Xaa₃₂₈ is Gln

Xaa₃₃₄ is Val

Xaa₃₄₂ is Ile

Xaa₃₇₇ is Thr

Xaa₃₈₁ is Glu

Xaa₃₈₅ is His

Xaa₃₈₇ is Ile

Xaa₃₉₃ is Val

Xaa₃₉₄ is Leu

Xaa₄₀₂ is Arg

Xaa₄₀₄ is Pro

Xaa₄₁₃ is Phe

Xaa₄₂₂ is Gly

Xaa₄₂₈ is Gly

Xaa₄₂₉ is Pro

Xaa₄₃₅ is GIn

Xaa₄₄₇ is Arg

Xaa₄₅₃ is Asn

Xaa₄₅₉ is Met, and

Xaa₄₈₅ is Asp.

Claim 55. (new) The isolated nucleic acid fragment of Claim 1 where

Xaa₁₀ is Phe or Leu

Xaa₁₆ is Leu

Xaa₂₃ is Thr

Xaa₂₅ is Lys

Xaa₃₉ is Lys

Xaa₄₈ is Leu

Xaa₆₀ is Leu

Xaa₇₃ is Leu

Xaa₇₄ is Tyr

Xaa₉₅ is Thr

Xaa₉₆ is His

Xaa₁₀₂ is Asn

Xaa₁₁₀ is lle

Xaa₁₁₂ is Arg

Xaa₁₁₇ is Ser

Xaa₁₁₈ is Ser

Xaa₁₂₁ is Met

Xaa₁₂₂ is Val

Xaa₁₂₄ is Phe

Xaa₁₂₉ is Lys

Xaa₁₄₇ is Lys

Xaa₁₅₉ is Phe

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Xaa₁₆₂ is Val

Xaa₁₆₆ is Gly Xaa₁₇₀ is Gln

Xaa₁₇₅ is Leu

Xaa₁₈₃ is Thr

Xaa₁₈₇ is Ile

Xaa₁₉₁ is Val

Xaa₂₀₉ is Phe

Xaa₂₁₉ is Trp

Xaa₂₂₃ is His

Xaa₂₅₃ is Glu

Xaa₂₅₉ is Lys

Xaa₂₆₃ is Val

Xaa₂₆₄ is Val

Xaa₂₆₈ is Val

Xaa₂₇₂ is Phe

Xaa₂₈₅ is Met

Xaa₂₉₂ is Asp

Xaa₂₉₃ is His

Xaa₂₉₄ is Ile

Xaa₃₀₁ is Phe

Xaa₃₀₆ is Thr

Xaa₃₁₁ is Val

Xaa₃₁₂ is Ala

Xaa₃₂₅ is Lys

Xaa₃₂₈ is Glu

Xaa₃₃₄ is Val

Xaa₃₄₂ is Arg

Xaa₃₇₇ is Thr

Xaa₃₈₁ is Glu

Xaa₃₈₅ is Cys

Xaa₃₈₇ is Thr

Xaa₃₉₃ is Ile

Xaa₃₉₄ is Leu

Xaa₄₀₂ is Arg

Xaa₄₀₄ is Pro

Xaa₄₁₃ is Phe

Xaa₄₂₂ is Gly

Xaa₄₂₈ is Arg

Xaa₄₂₉ is Leu

Xaa₄₃₅ is Arg

Xaa₄₄₇ is Arg

Xaa₄₅₃ is Asn

Xaa₄₅₉ is Met, and

Xaa₄₈₅ is Gly.

Claim 56. (new) The isolated nucleic acid fragment of Claim 1 where

Xaa₁₀ is Leu

Xaa₁₆ is Leu

Xaa₂₃ is Thr

Xaa₂₅ is Lys

Xaa₃₉ is Lys

Xaa₄₈ is Leu

Xaa₆₀ is Leu

Xaa₇₃ is Leu

Xaa₇₄ is Tyr

Xaa₉₅ is Thr

Xaa₉₆ is His

Xaa₁₀₂ is Asn

Xaa₁₁₀ is Ile

Xaa₁₁₂ is Arg

Xaa₁₁₇ is Ser

Xaa₁₁₈ is Ser

Xaa₁₂₁ is Met

Xaa₁₂₂ is Val

Xaa₁₂₄ is Phe

Xaa₁₂₉ is Lys

Xaa₁₄₇ is Lys

Xaa₁₅₉ is Phe

Xaa₁₆₂ is Val

Xaa₁₆₆ is Gly

Xaa₁₇₀ is Gln

Xaa₁₇₅ is Leu

Xaa₁₈₃ is Thr

Xaa₁₈₇ is Ile

Xaa₁₉₁ is Met

Xaa₂₀₉ is Phe

Xaa₂₁₉ is Trp

Xaa₂₂₃ is His

Xaa₂₅₃ is Glu

Xaa₂₅₉ is Lys

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Xaa₂₆₃ is Val

Xaa₂₆₄ is Val

Xaa₂₆₈ is Val

Xaa₂₇₂ is Phe

Xaa₂₈₅ is Met

Xaa₂₉₂ is Asp

Xaa₂₉₃ is His

Xaa₂₉₄ is Ile

Xaa₃₀₁ is Phe

Xaa₃₀₆ is Thr

Xaa₃₁₁ is Val

Xaa₃₁₂ is Ala

Xaa₃₂₅ is Lys

Xaa₃₂₈ is Glu

Xaa₃₃₄ is Ala

Xaa₃₄₂ is Arg

Xaa₃₇₇ is Thr

Xaa₃₈₁ is Glu

Xaa₃₈₅ is Tyr

Xaa₃₈₇ is Ile

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Xaa₃₉₃ is Ile

Xaa₃₉₄ is Leu

Xaa₄₀₂ is Arg

Xaa₄₀₄ is Pro

Xaa₄₁₃ is Phe

Xaa₄₂₂ is Gly

Xaa₄₂₈ is Arg

Xaa₄₂₉ is Pro

Xaa₄₃₅ is Gln

Xaa₄₄₇ is Arg

Xaa₄₅₃ is Asn

Xaa₄₅₉ is Met, and

Xaa₄₈₅ is Gly.

Claim 57. (new) The isolated nucleic acid fragment of Claim 1 where

Xaa₁₀ is Leu

Xaa₁₆ is Leu

Xaa₂₃ is Thr

Xaa₂₅ is Lys

Xaa₃₉ is Lys

Xaa₄₈ is Leu

Xaa₆₀ is Leu

Xaa₇₃ is Leu

Xaa₇₄ is Tyr

Xaa₉₅ is Thr

Xaa₉₆ is His

Xaa₁₀₂ is Asn

Xaa₁₁₀ is Ile

Xaa₁₁₂ is Arg

Xaa₁₁₇ is Ser

Xaa₁₁₈ is Ser

Xaa₁₂₁ is Met

Xaa₁₂₂ is Val

Xaa₁₂₄ is Phe

Xaa₁₂₉ is Lys

Xaa₁₄₇ is Lys

Xaa₁₅₉ is Phe

Xaa₁₆₂ is Ala

Xaa₁₆₆ is Gly

Xaa₁₇₀ is Gln

Xaa₁₇₅ is Leu

Xaa₁₈₃ is Thr

Xaa₁₈₇ is Ile

Xaa₁₉₁ is Met

Xaa₂₀₉ is Phe

Xaa₂₁₉ is Trp

Xaa₂₂₃ is His

Xaa₂₅₃ is Glu

Xaa₂₅₉ is Lys

Xaa₂₆₃ is Val

Xaa₂₆₄ is Val

Xaa₂₆₈ is Val

Xaa₂₇₂ is Phe

Xaa₂₈₅ is Met

Xaa₂₉₂ is Asp

Xaa₂₉₃ is His

Xaa₂₉₄ is lle

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Xaa₃₀₁ is Phe

Xaa₃₀₆ is Thr

Xaa₃₁₁ is Val

Xaa₃₁₂ is Ala

Xaa₃₂₅ is Lys

Xaa₃₂₈ is Glu

Xaa₃₃₄ is Val

Xaa₃₄₂ is Arg

Xaa₃₇₇ is Thr

Xaa₃₈₁ is Glu

Xaa₃₈₅ is Tyr

Xaa₃₈₇ is lle

Xaa₃₉₃ is lle

Xaa₃₉₄ is Leu

Xaa₄₀₂ is Arg

Xaa₄₀₄ is Pro

Xaa₄₁₃ is Phe

Xaa₄₂₂ is Gly

Xaa₄₂₈ is Arg

Xaa₄₂₉ is Pro

Xaa₄₃₅ is Gln

Xaa₄₄₇ is Arg

Xaa₄₅₃ is Asn

Xaa₄₅₉ is Met, and

Xaa₄₈₅ is Gly.

Claim 58. (new) The isolated nucleic acid fragment of Claim 1 where

Xaa₁₀ is Leu

Xaa₁₆ is Leu

Xaa₂₃ is Thr

Xaa₂₅ is Lys

Xaa₃₉ is Lys

Xaa₄₈ is Leu

Xaa₆₀ is Leu

Xaa₇₃ is Leu

Xaa₇₄ is Tyr

Xaa₉₅ is Thr Xaa₉₆ is His

Xaa₁₀₂ is Asn

Xaa₁₁₀ is Iler

Xaa₁₁₂ is Arg

Xaa₁₁₇ is Ser

Xaa₁₁₈ is Ser

Xaa₁₂₁ is Met

Xaa₁₂₂ is Val

Xaa₁₂₄ is Phe

Xaa₁₂₉ is Lys

Xaa₁₄₇ is Lys

Xaa₁₅₉ is Phe

Xaa₁₆₂ is Val

Xaa₁₆₆ is Gly

Xaa₁₇₀ is Gln

Xaa₁₇₅ is Leu

Xaa₁₈₃ is Thr

Xaa₁₈₇ is lle

Xaa₁₉₁ is Met

Xaa₂₀₉ is Phe

Xaa₂₁₉ is Trp

Xaa₂₂₃ is His

Xaa₂₅₃ is Glu

Xaa₂₅₉ is Lys

Xaa₂₆₃ is Val

Xaa₂₆₄ is Val

Xaa₂₆₈ is Val

Xaa₂₇₂ is Phe

Xaa₂₈₅ is Thr

Xaa₂₉₂ is Asp

Xaa₂₉₃ is His

Xaa₂₉₄ is lle

Xaa₃₀₁ is Phe

Xaa₃₀₆ is Thr

Xaa₃₁₁ is Val

Xaa₃₁₂ is Ala

Xaa₃₂₅ is Lys

Xaa₃₂₈ is Glu

Xaa₃₃₄ is Val

Xaa₃₄₂ is Arg

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Xaa₃₇₇ is Thr

Xaa₃₈₁ is Glu

Xaa₃₈₅ is Tyr

Xaa₃₈₇ is lle

Xaa₃₉₃ is lle

Xaa₃₉₄ is Leu

Xaa₄₀₂ is Arg

Xaa₄₀₄ is Pro

Xaa₄₁₃ is Phe

Xaa₄₂₂ is Gly

Xaa₄₂₈ is Arg

Xaa₄₂₉ is Pro

Xaa₄₃₅ is Gln

Xaa₄₄₇ is Arg

Xaa₄₅₃ is Asn

Xaa₄₅₉ is Met, and

Xaa₄₈₅ is Gly.

Claim 59. (new) The isolated nucleic acid fragment of Claim 1 where

Xaa₁₀ is Leu

Xaa₁₆ is Leu

Xaa₂₃ is Thr

Xaa₂₅ is Lys

Xaa₃₉ is Lys

Xaa₄₈ is Leu

Xaa₆₀ is Leu

Xaa₇₃ is Leu

Xaa₇₄ is Tyr

Xaa₉₅ is Thr

Vaa is His

Xaa₉₆ is His

Xaa₁₀₂ is Asn

Xaa₁₁₀ is lle

Xaa₁₁₂ is Arg

Xaa₁₁₇ is Ser

Xaa₁₁₈ is Ser

Xaa₁₂₁ is Met

Xaa₁₂₂ is Val

Xaa₁₂₄ is Phe

Xaa₁₂₉ is Lys

Xaa₁₄₇ is Lys

Xaa₁₅₉ is Phe

Xaa₁₆₂ is Val

Xaa₁₆₆ is Gly

Xaa₁₇₀ is Gln

Xaa₁₇₅ is Leu

Xaa₁₈₃ is Thr

Xaa₁₈₇ is Ile

Xaa₁₉₁ is Met

Xaa₂₀₉ is Phe

Xaa₂₁₉ is Trp

Xaa₂₂₃ is His

Xaa₂₅₃ is Glu

Xaa₂₅₉ is Lys

Xaa₂₆₃ is Val

Xaa₂₆₄ is Val

Xaa₂₆₈ is Val

Xaa₂₇₂ is Phe

Xaa₂₈₅ is Met

Xaa₂₉₂ is Asp

Xaa₂₉₃ is His

Xaa₂₉₄ is Ile

Xaa₃₀₁ is Phe

Xaa₃₀₆ is Thr

Xaa₃₁₁ is Glu

Xaa₃₁₂ is Ala

Xaa₃₂₅ is Lys

Xaa₃₂₈ is Glu

Xaa₃₃₄ is Val

Xaa₃₄₂ is Arg

Xaa₃₇₇ is Thr

Xaa₃₈₁ is Glu

Xaa₃₈₅ is Tyr

Xaa₃₈₇ is Ile

Xaa₃₉₃ is Ile

Xaa₃₉₄ is Leu

Xaa₄₀₂ is Arg

Xaa₄₀₄ is Pro

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Xaa₄₁₃ is Phe

Xaa₄₂₂ is Gly

Xaa₄₂₈ is Arg

Xaa₄₂₉ is Pro

Xaa₄₃₅ is Gln

Xaa₄₄₇ is Arg

Xaa₄₅₃ is Asn

Xaa₄₅₉ is Met, and

Xaa₄₈₅ is Gly.

Claim 60. (new) The isolated nucleic acid fragment of Claim 1 where

Xaa₁₀ is Leu

Xaa₁₆ is Leu

Xaa₂₃ is Thr

Xaa₂₅ is Lys

Xaa₃₉ is Lys

Xaa₄₈ is Leu

Xaa₆₀ is Leu

Xaa₇₃ is Leu

Xaa₇₄ is Tyr

Xaa₉₅ is Thr

Xaa₉₆ is His

Xaa₁₀₂ is Asn

Xaa₁₁₀ is Ile

Xaa₁₁₂ is Arg

Xaa₁₁₇ is Ser

Xaa₁₁₈ is Ser

Xaa₁₂₁ is Met

Xaa₁₂₂ is Val

Xaa₁₂₄ is lle

Xaa₁₂₉ is Lys

Xaa₁₄₇ is Lys

Xaa₁₅₉ is Phe

Xaa₁₆₂ is Val

Xaa₁₆₆ is Gly

Xaa₁₇₀ is Gln

Xaa₁₇₅ is Leu

Xaa₁₈₃ is Thr

Xaa₁₈₇ is Ile

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Xaa₁₉₁ is Met

Xaa₂₀₉ is Phe

Xaa₂₁₉ is Trp

Xaa₂₂₃ is His

Xaa₂₅₃ is Glu

Xaa₂₅₉ is Lys

Xaa₂₆₃ is Val

Xaa₂₆₄ is Asp

Xaa₂₆₈ is Val

Xaa₂₇₂ is Phe

Xaa₂₈₅ is Thr

Xaa₂₉₂ is Asp

Xaa₂₉₃ is His

Xaa₂₉₄ is Ile

Xaa₃₀₁ is Phe

Xaa₃₀₆ is Thr

Xaa₃₁₁ is Val

Xaa₃₁₂ is Ala

Xaa₃₂₅ is Lys

Xaa₃₂₈ is Glu

Xaa₃₃₄ is Val

Xaa₃₄₂ is Arg

Xaa₃₇₇ is Thr

Xaa₃₈₁ is Glu

Xaa₃₈₅ is Tyr

Xaa₃₈₇ is lle

Xaa₃₉₃ is lle

Xaa₃₉₄ is Leu

Xaa₄₀₂ is Arg

Xaa₄₀₄ is Pro

Xaa₄₁₃ is Phe

Xaa₄₂₂ is Gly

Xaa₄₂₈ is Arg

Xaa₄₂₉ is Pro

Xaa₄₃₅ is GIn

Xaa₄₄₇ is Arg

Xaa₄₅₃ is Asn

Xaa₄₅₉ is Met

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Xaa₄₈₅ is Gly.

Claim 61. (new) The isolated nucleic acid fragment of Claim 1 where

Xaa₁₀ is Leu

Xaa₁₆ is Leu

Xaa₂₃ is Thr

Xaa₂₅ is Lys

Xaa₃₉ is Lys

Xaa₄₈ is Leu

Xaa₆₀ is Leu

Xaa₇₃ is Leu

Xaa₇₄ is Tyr

Xaa₉₅ is Thr

Xaa₉₆ is His

Xaa₁₀₂ is Asn

Xaa₁₁₀ is Ile

Xaa₁₁₂ is Arg

Xaa₁₁₇ is Ser

Xaa₁₁₈ is Ser

Xaa₁₂₁ is Met

Xaa₁₂₂ is Val

Xaa₁₂₄ is Phe

Xaa₁₂₉ is Lys

Xaa₁₄₇ is Lys

Xaa₁₅₉ is Phe

Xaa₁₆₂ is Val

Xaa₁₆₆ is Gly

Xaa₁₇₀ is Gln

Xaa₁₇₅ is Leu

Xaa₁₈₃ is Thr

Xaa₁₈₇ is lle

Xaa₁₉₁ is Met

Xaa₂₀₉ is Phe

Xaa₂₁₉ is Trp

Xaa₂₂₃ is His

Xaa₂₅₃ is Glu

Xaa₂₅₉ is Lys

Xaa₂₆₃ is Val

Xaa₂₆₄ is Val

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Xaa₂₆₈ is Val

Xaa₂₇₂ is Phe

 Xaa_{285} is Thr

Xaa₂₉₂ is Asp

Xaa₂₉₃ is His

Xaa₂₉₄ is Ile

Xaa₃₀₁ is Phe

Xaa₃₀₆ is Thr

Xaa₃₁₁ is Va

Xaa₃₁₂ is Ala

Xaa₃₂₅ is Lys

Xaa₃₂₈ is Glu

Xaa₃₃₄ is Val

Xaa₃₄₂ is Arg

Xaa₃₇₇ is Thr

Xaa₃₈₁ is Glu

Xaa₃₈₅ is Tyr

Xaa₃₈₇ is lle

Xaa₃₉₃ is lle

Xaa₃₉₄ is Leu

Xaa₄₀₂ is Arg

Xaa₄₀₄ is Pro

Xaa₄₁₃ is Phe

Xaa₄₂₂ is Gly

Xaa₄₂₈ is Arg

Xaa₄₂₉ is Pro

Xaa₄₃₅ is GIn

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Xaa₄₄₇ is Arg

Xaa₄₅₃ is Asn

Xaa₄₅₉ is Met, and

Xaa₄₈₅ is Gly.

Claim 62. (new) The isolated nucleic acid fragment of Claim 1 where

Xaa₁₀ is Phe

Xaa₁₆ is Leu

Xaa₂₃ is Ser

Xaa₂₅ is Lys

Xaa₃₉ is Lys

Xaa₄₈ is Leu

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Xaa₆₀ is Leu

Xaa₇₃ is Leu

Xaa₇₄ is Ser

Xaa₉₅ is Ala

Xaa₉₆ is His

Xaa₁₀₂ is Ser

Xaa₁₁₀ is Val

Xaa₁₁₂ is Arg

Xaa₁₁₇ is Asn

Xaa₁₁₈ is Ser

Xaa₁₂₁ is Met

Xaa₁₂₂ is Val

Xaa₁₂₄ is Phe

Xaa₁₂₉ is Lys

Xaa₁₄₇ is Glu

Xaa₁₅₉ is Phe

Xaa₁₆₂ is Val

Xaa₁₆₆ is Ser

Xaa₁₇₀ is Gln

Xaa₁₇₅ is Val

Xaa₁₈₃ is Thr

Xaa₁₈₇ is Ile

Xaa₁₉₁ is Met

Xaa₂₀₉ is Phe

Xaa₂₁₉ is Trp

Xaa₂₂₃ is Tyr

Xaa₂₅₃ is Glu

Xaa₂₅₉ is Lys

Xaa₂₆₃ is Val

Xaa₂₆₄ is Val

Xaa₂₆₈ is Ala

Xaa₂₇₂ is Phe

Xaa₂₈₅ is Met

Xaa₂₉₂ is Glu

Xaa₂₉₃ is Gln

Xaa₂₉₄ is lle

Xaa₃₀₁ is Phe

Xaa₃₀₆ is Thr

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Xaa₃₁₁ is Val

Xaa₃₁₂ is Ala

Xaa₃₂₅ is Arg

Xaa₃₂₈ is GIn

Xaa₃₃₄ is Val

Xaa₃₄₂ is Arg

Xaa₃₇₇ is Thr

Xaa₃₈₁ is Glu

Xaa₃₈₅ is Tyr

Xaa₃₈₇ is lle

Xaa₃₉₃ is Val

Xaa₃₉₄ is Leu

Xaa₄₀₂ is Lys

Xaa₄₀₄ is Pro

Xaa₄₁₃ is Phe

Xaa₄₂₂ is Gly

Xaa₄₂₈ is Gly

Xaa₄₂₉ is Pro

Xaa₄₃₅ is Gln

Xaa₄₄₇ is Arg

Xaa₄₅₃ is Asn

Xaa₄₅₉ is Met, and

Xaa₄₈₅ is Asp.

Claim 63. (new) The isolated nucleic acid fragment of Claim 1 where

Xaa₁₀ is Phe or Leu

Xaa₁₆ is Ser

Xaa₂₃ is Ser

Xaa₂₅ is Ile

Xaa₃₉ is Arg

Xaa₄₈ is Leu

Xaa₆₀ is Pro

Xaa₇₃ is Leu

Xaa₇₄ is Ser

Xaa₉₅ is Thr

Xaa₉₆ is His

Xaa₁₀₂ is Asn

Xaa₁₁₀ is Ile

Xaa₁₁₂ is His

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Xaa₁₁₇ is Asn

Xaa₁₁₈ is Ser

Xaa₁₂₁ is Met

Xaa₁₂₂ is Val

Xaa₁₂₄ is Phe

Xaa₁₂₉ is Lys

Xaa₁₄₇ is Lys

Xaa₁₅₉ is Phe

Xaa₁₆₂ is Val

Xaa₁₆₆ is Ser

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Xaa₁₇₀ is Gln

Xaa₁₇₅ is Val

Xaa₁₈₃ is Thr

Xaa₁₈₇ is Ile

Xaa₁₉₁ is Met

Xaa₂₀₉ is Phe

Xaa₂₁₉ is Trp

Xaa₂₂₃ is Tyr

Xaa₂₅₃ is Glu

Xaa₂₅₉ is Lys

Xaa₂₆₃ is Val

Xaa₂₆₄ is Val

Xaa₂₆₈ is Ala

Xaa₂₇₂ is Phe

Xaa₂₈₅ is Met

Xaa₂₉₂ is Glu

Xaa₂₉₃ is Gln

Xaa₂₉₄ is Ile

Xaa₃₀₁ is Phe

Xaa₃₀₆ is Thr

Xaa₃₁₁ is Val

Xaa₃₁₂ is Val

Xaa₃₂₅ is Arg

Xaa₃₂₈ is Gln

Xaa₃₃₄ is Val

Xaa₃₄₂ is Arg

Xaa₃₇₇ is Thr

Xaa₃₈₁ is Glu

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Xaa₃₈₅ is Tyr

Xaa₃₈₇ is lle

Xaa₃₉₃ is Val

Xaa₃₉₄ is Leu

Xaa₄₀₂ is Arg

Xaa₄₀₄ is Pro

Xaa₄₁₃ is Ser

Xaa₄₂₂ is Gly

Xaa₄₂₈ is Gly

Xaa₄₂₉ is Pro

Xaa₄₃₅ is Gln

Xaa₄₄₇ is Arg

Xaa₄₅₃ is Ser

Xaa₄₅₉ is Met, and

Xaa₄₈₅ is Asp.

Claim 64. (new) The isolated nucleic acid fragment of Claim 1 where

Xaa₁₀ is Phe or Leu

Xaa₁₆ is Leu

Xaa₂₃ is Thr

Xaa₂₅ is Lys

Xaa₃₉ is Lys

Xaa₄₈ is Leu

Xaa₆₀ is Leu

Xaa₇₃ is Leu

Xaa₇₄ is Tyr

Xaa₉₅ is Thr

Xaa₉₆ is His

Xaa₁₀₂ is Asn

Xaa₁₁₀ is Ile

Xaa₁₁₂ is Arg

Xaa₁₁₇ is Ser

Xaa₁₁₈ is Ser

Xaa₁₂₁ is Met

Xaa₁₂₂ is Val

Xaa₁₂₄ is Phe

Xaa₁₂₉ is Lys

Xaa₁₄₇ is Lys

Xaa₁₅₉ is Phe

Xaa₁₆₂ is Val

Xaa₁₆₆ is Ser

Xaa₁₇₀ is Gln

Xaa₁₇₅ is Val

Xaa₁₈₃ is Thr

Xaa₁₈₇ is Ile

Xaa₁₉₁ is Met

Xaa₂₀₉ is Phe

Xaa₂₁₉ is Trp

Xaa₂₂₃ is Tyr

Xaa₂₅₃ is Glu

Xaa₂₅₉ is Lys

Xaa₂₆₃ is Val

Xaa₂₆₄ is Val

Xaa₂₆₈ is Ala

Xaa₂₇₂ is Phe

Xaa₂₈₅ is Met

Xaa₂₉₂ is Glu

Xaa₂₉₃ is Gln

Xaa₂₉₄ is Ile

Xaa₃₀₁ is Phe

Xaa₃₀₆ is Thr

Xaa₃₁₁ is Val

Xaa₃₁₂ is Val

Xaa₃₂₅ is Lys

Xaa₃₂₈ is Gln

Xaa₃₃₄ is Ala

Xaa₃₄₂ is Arg

..

Xaa₃₇₇ is Thr Xaa₃₈₁ is Gly

Xaa₃₈₅ is Tyr

Xaa₃₈₇ is Ile

Xaa₃₉₃ is Val

Xaa₃₉₄ is Leu

Xaa₄₀₂ is Arg Xaa₄₀₄ is Pro

Xaa₄₁₃ is Phe

Xaa₄₂₂ is Gly

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Xaa₄₂₈ is Gly

Xaa₄₂₉ is Pro

Xaa₄₃₅ is GIn

Xaa₄₄₇ is Arg

Xaa₄₅₃ is Asn

Xaa₄₅₉ is Met, and

Xaa₄₈₅ is Asp.

Claim 65. (new) The isolated nucleic acid fragment of Claim 1 where

Xaa₁₀ is Phe or Leu

Xaa₁₆ is Leu

Xaa₂₃ is Thr

Xaa₂₅ is Lys

Xaa₃₉ is Lys

Xaa₄₈ is Leu

Xaa₆₀ is Leu

Xaa₇₃ is Leu

Xaa₇₄ is Tyr

Xaa₉₅ is Thr

Xaa₉₆ is His

Xaa₁₀₂ is Asn

Xaa₁₁₀ is Ile

Xaa₁₁₂ is Arg

Xaa₁₁₇ is Ser

Xaa₁₁₈ is Ser

Xaa₁₂₁ is Met

Xaa₁₂₂ is Val

Xaa₁₂₄ is Phe

Xaa₁₂₉ is Lys

Xaa₁₄₇ is Lys

Xaa₁₅₉ is Phe

Xaa₁₆₂ is Val

Xaa₁₆₆ is Gly

Xaa₁₇₀ is Gln

7133170 15 5111

Xaa₁₇₅ is Leu Xaa₁₈₃ is Thr

V-- !- !!-

Xaa₁₈₇ is Ile

Xaa₁₉₁ is Met

Xaa₂₀₉ is Phe

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Xaa₂₁₉ is Trp

Xaa₂₂₃ is His

Xaa₂₅₃ is Glu

Xaa₂₅₉ is Lys

Xaa₂₆₃ is Asp

Xaa₂₆₄ is Val

Xaa₂₆₈ is Val

Xaa₂₇₂ is Phe

Xaa₂₈₅ is Met

Xaa₂₉₂ is Asp

Xaa₂₉₃ is His

Xaa₂₉₄ is Ile

Xaa₃₀₁ is Phe

Xaa₃₀₆ is Thr

Xaa₃₁₁ is Val

Xaa₃₁₂ is Ala

Xaa₃₂₅ is Lys

Xaa₃₂₈ is Glu

Xaa₃₃₄ is Val

Xaa₃₄₂ is Arg

Xaa₃₇₇ is lle

Xaa₃₈₁ is Glu

Xaa₃₈₅ is Tyr

Xaa₃₈₇ is lle

Xaa₃₉₃ is lle

Xaa₃₉₄ is Leu

Xaa₄₀₂ is Arg

Xaa₄₀₄ is Pro

Xaa₄₁₃ is Phe

Xaa₄₂₂ is Gly

Xaa₄₂₈ is Arg

Xaa₄₂₉ is Leu

Xaa₄₃₅ is Gln

Xaa₄₄₇ is Arg

Xaa₄₅₃ is Asn

Xaa₄₅₉ is Met, and

Xaa₄₈₅ is Gly.

Claim 66. (new) The isolated nucleic acid fragment of claim 1 where

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Xaa₁₀ is Phe or Leu

Xaa₁₆ is Leu

Xaa₂₃ is Thr

Xaa₂₅ is Lys

Xaa₃₉ is Lys

Xaa₄₈ is Leu

Xaa₆₀ is Leu

Xaa₇₃ is Leu

Xaa₇₄ is Tyr

Xaa₉₅ is Thr

Xaa₉₆ is His

Xaa₁₀₂ is Asn

Xaa₁₁₀ is lle

Xaa₁₁₂ is Arg

Xaa₁₁₇ is Ser

Xaa₁₁₈ is Ser

Xaa₁₂₁ is Arg

Xaa₁₂₂ is Val

Xaa₁₂₄ is Phe

Xaa₁₂₉ is Lys

Xaa₁₄₇ is Lys

Xaa₁₅₉ is Phe

Xaa₁₆₂ is Val

Xaa₁₆₆ is Gly

Xaa₁₇₀ is Gln

Xaa₁₇₅ is Leu

Xaa₁₈₃ is Thr

Xaa₁₈₇ is lle

Xaa₁₉₁ is Metl

Xaa₂₀₉ is Phe

Xaa₂₁₉ is Trp

Xaa₂₂₃ is His

Xaa₂₅₃ is Glu

Xaa₂₅₉ is Lys

Xaa₂₆₃ is Val

Xaa₂₆₄ is Val

Xaa₂₆₈ is Val

Xaa₂₇₂ is Leu

Xaa₂₈₅ is Met

Xaa₂₉₂ is Asp

Xaa₂₉₃ is His

Xaa₂₉₄ is lle

Xaa₃₀₁ is Phe

Xaa₃₀₆ is Thr

Xaa₃₁₁ is Val

Xaa312 is Ala

Xaa₃₂₅ is Lys

Xaa₃₂₈ is Glu

Xaa₃₃₄ is Val

Xaa₃₄₂ is Arg

Xaa₃₇₇ is Thr

Xaa₃₈₁ is Glu

Xaa₃₈₅ is Tyr

Xaa₃₈₇ is Ile

Xaa₃₉₃ is Ile

Xaa₃₉₄ is Leu

Xaa₄₀₂ is Arg

Xaa₄₀₄ is Pro

Xaa₄₁₃ is Phe

Xaa₄₂₂ is Glu

Xaa₄₂₈ is Arg

Xaa₄₂₉ is Pro

Xaa₄₃₅ is Gln

Xaa₄₄₇ is Arg

Xaa₄₅₃ is

Xaa₄₅₉ is Met, and

Xaa₄₈₅ is Gly.

Claim 67. (new) The isolated nucleic acid fragment of Claim 1 where

Xaa₁₀ is Phe or Leu

Xaa₁₆ is Leu

Xaa₂₃ is Thr

Xaa₂₅ is Lys

Xaa₃₉ is Lys

Xaa₄₈ is Leu

Xaa₆₀ is Leu

Xaa₇₃ is Leu

Xaa₇₄ is Tyr

Xaa₉₅ is Thr

Xaa₉₆ is His

Xaa₁₀₂ is Asn

Xaa₁₁₀ is Ile

Xaa₁₁₂ is Arg

Xaa₁₁₇ is Ser

Xaa₁₁₈ is Ser

Xaa₁₂₁ is Met

Xaa₁₂₂ is Ala

Xaa₁₂₄ is Phe

Xaa₁₂₉ is Lys

Xaa₁₄₇ is Lys

Xaa₁₅₉ is Phe

Xaa₁₆₂ is Val

Xaa₁₆₆ is Gly

Xaa₁₇₀ is Gln

Xaa₁₇₅ is Leu

Xaa₁₈₃ is Thr

Xaa₁₈₇ is Thr

Xaa₁₉₁ is Met

Xaa₂₀₉ is Phe

Xaa₂₁₉ is Arg

Xaa₂₂₃ is His

Xaa₂₅₃ is Glu

Xaa₂₅₉ is Lys

Xaa₂₆₃ is Val

Xaa₂₆₄ is Val

Xaa₂₆₈ is Val

Xaa₂₇₂ is Phe

Xaa₂₈₅ is Thr

Xaa₂₉₂ is Asp

Xaa₂₉₃ is His

Xaa₂₉₄ is Ile

Xaa₃₀₁ is Phe

Xaa₃₀₆ is Thr

Xaa₃₁₁ is Val

Xaa₃₁₂ is Ala

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Xaa₃₂₅ is Lys

Xaa₃₂₈ is Glu

Xaa₃₃₄ is Val

Xaa₃₄₂ is Arg

Xaa₃₇₇ is Thr

Xaa₃₈₁ is Glu

Xaa₃₈₅ is Tyr

Xaa₃₈₇ is Ile

Xaa₃₉₃ is Ile

Xaa₃₉₄ is Leu

Xaa₄₀₂ is Arg

Xaa₄₀₄ is Ser

Y '- Di-

Xaa₄₁₃ is Phe

Xaa₄₂₂ is Gly Xaa₄₂₈ is Arg

V-- '- D--

Xaa₄₂₉ is Pro

Xaa₄₃₅ is Gln

Xaa₄₄₇ is Arg

Xaa₄₅₃ is Asn

Xaa₄₅₉ is Met, and

Xaa₄₈₅ is Gly.

Claim 68. (new) The isolated nucleic acid fragment of Claim 1 where

Xaa₁₀ is Phe or Leu

Xaa₁₆ is Leu

Xaa₂₃ is Thr

Xaa₂₅ is Lys

Xaa₃₉ is Lys

Xaa₄₈ is Leu

Xaa₆₀ is Leu

Xaa₇₃ is Leu

Xaa₇₄ is Tyr

Xaa₉₅ is Thr

Xaa₉₆ is His

Xaa₁₀₂ is Asn

102

Xaa₁₁₀ is Ile

Xaa₁₁₂ is Arg

Xaa₁₁₇ is Ser

Xaa₁₁₈ is Ser

Xaa₁₂₁ is Met

Xaa₁₂₂ is Val

Xaa₁₂₄ is Phe

Xaa₁₂₉ is Lys

Xaa₁₄₇ is Lys

Xaa₁₅₉ is Leu

Xaa₁₆₂ is Val

Xaa₁₆₆ is Gly

Xaa₁₇₀ is Gln

Xaa₁₇₅ is Leu

Xaa₁₈₃ is Thr

Xaa₁₈₇ is Ile

Xaa₁₉₁ is Met

Xaa₂₀₉ is Phe

Xaa₂₁₉ is Trp

Xaa₂₂₃ is His

Xaa₂₅₃ is Glu

Xaa₂₅₉ is Lys

Xaa₂₆₃ is Val

Xaa₂₆₄ is Ile

Xaa₂₆₈ is Val

Xaa₂₇₂ is Phe

Xaa₂₈₅ is Thr

Xaa₂₉₂ is Asp

Xaa₂₉₃ is His

Xaa₂₉₄ is Ile

Xaa₃₀₁ is Phe

Xaa₃₀₆ is Thr

Xaa₃₁₁ is Val

Xaa₃₁₂ is Ala

Xaa₃₂₅ is Lys

Xaa₃₂₈ is Glu

Xaa₃₃₄ is Val

Xaa₃₄₂ is Arg

Xaa₃₇₇ is Thr

Xaa₃₈₁ is Glu

Xaa₃₈₅ is Tyr

Xaa₃₈₇ is Ile

Xaa₃₉₃ is Ile

Xaa₃₉₄ is Leu

Xaa₄₀₂ is Arg

Xaa₄₀₄ is Pro

Xaa₄₁₃ is Phe

Xaa₄₂₂ is Gly

Xaa₄₂₈ is Arg

Xaa₄₂₉ is Pro

Xaa₄₃₅ is Gln

Xaa₄₄₇ is Arg

Xaa₄₅₃ is Asn

Xaa₄₅₉ is Met, and

Xaa₄₈₅ is Gly.

Claim 69. (new) The isolated nucleic acid fragment of Claim 1 where

Xaa₁₀ is Phe or Leu

Xaa₁₆ is Leu

Xaa₂₃ is Thr

Xaa₂₅ is Lys

Xaa₃₉ is Lys

Xaa₄₈ is Leu

Xaa₆₀ is Leu

Xaa₇₃ is Leu or His

Xaa₇₄ is Ser or Tyr

Xaa₉₅ is Ala or Thr

Xaa₉₆ is Asn or His

Xaa₁₀₂ is Asn or Ser

Xaa₁₁₀ is Ile, Val, or Thr

Xaa₁₁₂ is Arg or His

Xaa₁₁₇ is Asn or Ser

Xaa₁₁₈ is Ser or Leu

Xaa₁₂₁ is Met or Arg

Xaa₁₂₂ is Ala or Val

Xaa₁₂₄ is Phe or Ile

Xaa₁₂₉ is Lys or Arg

Xaa₁₄₇ is Lys or Glu

Xaa₁₅₉ is Leu or Phe

Xaa₁₆₂ is Ala or Val

Xaa₁₆₆ is Ser or Gly

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Xaa₁₇₀ is Gln or Arg

Xaa₁₇₅ is Val or Leu

Xaa₁₈₃ is Ala or Thr

Xaa₁₈₇ is Thr or Ile

Xaa₁₉₁ is Met or Val

Xaa₂₀₉ is Phe or Tyr

Xaa₂₁₉ is Arg or Trp

Xaa₂₂₃ is Tyr or His

Xaa₂₅₃ is Gly or Glu

Xaa₂₅₉ is Lys or Glu

Xaa₂₆₃ is Val or Asp

Xaa₂₆₄ is Val, Asp, or lle

Xaa₂₆₈ is Ala or Val

Xaa₂₇₂ is Phe or Leu

Xaa₂₈₅ is Thr or Met

Xaa₂₉₂ is Glu or Asp

Xaa₂₉₃ is Gln or His

Xaa₂₉₄ is Thr, or lle

Xaa₃₀₁ is Phe or Leu

Xaa₃₀₆ is Thr or Ile

Xaa₃₁₁ is Val or Glu

Xaa312 is Val or Ala

Xaa₃₂₅ is Arg or Lys

Xaa328 is Gln or Glu

Xaa₃₃₄ is Val or Ala

Xaa₃₄₂ is Arg or lle

Xaa377 is Thr or Ile

Xaa₃₈₁ is Glu or Gly

Xaa₃₈₅ is Tyr, His, or Cys

Xaa₃₈₇ is lle or Thr

Xaa₃₉₃ is Val or lie

Xaa₃₉₄ is Leu or Pro

Xaa₄₀₂ is Arg or Lys

Xaa₄₀₄ is Ser or Pro

Xaa₄₁₃ is Ser or Phe

Xaa₄₂₂ is Glu or Gly

Xaa₄₂₈ is Gly or Arg

Xaa₄₂₉ is Pro or Leu

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Xaa₄₃₅ is Gln or Arg Xaa₄₄₇ is Arg or Gly

Xaa₄₅₃ is Asn, Ser, or lle Xaa₄₅₉ is Met or Thr, and

Xaa₄₈₅ is Asp or Gly.